

SEQUENCE LISTING

<110> Robert G. Korneluk et al.

<120> METHODS AND COMPOUNDS FOR MODULATING
MALE FERTILITY

<130> 07891/018002

<140> 09/239,867

<141> 1999-01-29

<150> 60/073,001

<151> 1998-01-29

<160> 10

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1559

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(1559)

<223> n = A,T,C or G

<400> 1

gagacggtgg	acaagtccta	tattcaagag	aagataactt	tgaacagttt	cgaaggatct	60
aaaacgtatg	tgtctgcaga	catcaatnag	gatgaagaat	tagtanaaga	gattaataga	120
tcaaaaacgt	ttgctggctt	tgcaggtggt	gggcctgcct	gggcatcggc	gcgttggagg	180
agacgccctg	gggggcctta	gctgccctga	agcggtagac	aggtggcaac	gtgggggctc	240
aggagttagc	aaacacaaga	aagcagcgcc	gaattgcagg	tttatccgca	gcttttattt	300
tgaagacagt	gccacgaaac	ctgcaaattc	tgggtgtcca	aatagtcaat	accaagttga	360
aaaccatctg	ggagaggaaa	agcgttgtgc	tttagacagg	ccgtatgaga	ctcgtgcaga	420
ccggcttttg	agagctggac	aggtggtgga	tagatcagac	tccatacacc	cgaggagccc	480
cgccatgcat	agtgaagaag	ctagataaca	gtcgtttcac	aactggccag	cctctgcccc	540
cttgaccccg	agagagctgg	ccagtgtctg	gctgtactac	acaggcactg	atgaccaagt	600
gcagtgtctc	tgttgtggcg	gaaaactgaa	aaactgggaa	cctggtgatc	gtgcctggtc	660
agaacacagg	agacattttc	ctaattgctt	ctttattttg	ggccacaacg	ttaatattcg	720
aggtgaatct	gatgttgcca	gttctgatag	gaattttctc	aattcaacaa	gttctccaag	780
gaatccatcc	atgacgggtt	atgaagcccg	gctcattact	tttgggacat	ggatgtactc	840
cgtcaacaaa	gagcagcttg	caagagctgg	attttatgct	ataggtcaag	aggataaagt	900
acagtgtctt	cactgtggag	gagggctagc	caactggaag	ccaagggaag	atccttggga	960
acagcatgct	aaatggtatc	caggttgcaa	atatctgcta	gaagagaagg	gacatgaata	1020
tataaacaac	attcatttaa	cccgttcact	tgagggagct	ctggtacaaa	ctaccaagaa	1080
aacaccatca	ctaactaaaa	gaatcagtga	taccatcttc	cctaataccta	tgctacaaga	1140
agctatacga	atgggatttg	atttcaagga	cgtaagaaa	ataatggagg	aaagaattca	1200

```

aacatctggg agcaactata aaacgcttga ggttcttggt gcagatctag tgagcgctca 1260
gaaagacact acagaaaatg aattgaatca gacttcattg cagagagaaa tcagccctga 1320
agagccgcta aggcgtctgc aagaggagaa gctttgtaaa atctgcatgg acagatatat 1380
cgctgttggt tttattcctt gtggacatct ggtcacttgt aaacaatgtg ctgaagcagt 1440
tgacagatgt cccatgtgca gcgcggttat tgatttcaag caaagagttt ttatgtctta 1500
atgtaactct acagtgggtg tgctatgttc ttattaccct gattaaatgt gtgatgtga 1559

```

```

<210> 2
<211> 236
<212> PRT
<213> Homo sapiens

```

```

<400> 2
Met Thr Gly Tyr Glu Ala Arg Leu Ile Thr Phe Gly Thr Trp Met Tyr
  1          5          10          15
Ser Val Asn Lys Glu Gln Leu Ala Arg Ala Gly Phe Tyr Ala Ile Gly
      20          25          30
Gln Glu Asp Lys Val Gln Cys Phe His Cys Gly Gly Gly Leu Ala Asn
      35          40          45
Trp Lys Pro Lys Glu Asp Pro Trp Glu Gln His Ala Lys Trp Tyr Pro
      50          55          60
Gly Cys Lys Tyr Leu Leu Glu Glu Lys Gly His Glu Tyr Ile Asn Asn
      65          70          75          80
Ile His Leu Thr Arg Ser Leu Glu Gly Ala Leu Val Gln Thr Thr Lys
      85          90          95
Lys Thr Pro Ser Leu Thr Lys Arg Ile Ser Asp Thr Ile Phe Pro Asn
      100          105          110
Pro Met Leu Gln Glu Ala Ile Arg Met Gly Phe Asp Phe Lys Asp Val
      115          120          125
Lys Lys Ile Met Glu Glu Arg Ile Gln Thr Ser Gly Ser Asn Tyr Lys
      130          135          140
Thr Leu Glu Val Leu Val Ala Asp Leu Val Ser Ala Gln Lys Asp Thr
      145          150          155          160
Thr Glu Asn Glu Leu Asn Gln Thr Ser Leu Gln Arg Glu Ile Ser Pro
      165          170          175
Glu Glu Pro Leu Arg Arg Leu Gln Glu Glu Lys Leu Cys Lys Ile Cys
      180          185          190
Met Asp Arg Tyr Ile Ala Val Val Phe Ile Pro Cys Gly His Leu Val
      195          200          205
Thr Cys Lys Gln Cys Ala Glu Ala Val Asp Arg Cys Pro Met Cys Ser
      210          215          220
Ala Val Ile Asp Phe Lys Gln Arg Val Phe Met Ser
      225          230          235

```

```

<210> 3
<211> 1588
<212> DNA
<213> Homo sapiens

```

```

<400> 3
gaaaagggtg acaagtccta ttttcaagag aagatgactt ttaacagttt tgaaggatct 60

```

aaaacttg	tacctgcaga	catcaataag	gaagaagaat	ttgtagaaga	gtttaataga	120
ttaaaaactt	ttgctaattt	tccaagtgg	agtcctgttt	cagcatcaac	actggcacga	180
gcagggtttc	tttatactgg	tgaaggagat	accgtgcggt	gctttagttg	tcatgcagct	240
gtagatagat	ggcaatatgg	agactcagca	gttgggaagac	acaggaaagt	atcccccatt	300
tgcagattta	tcaacggcct	ttatcttgaa	aatagtgcc	cgcagtctac	aaattctggt	360
atccagaatg	gtcagtacaa	agttgaaaac	tatctgggaa	gcagagatca	ttttgcctta	420
gacaggccat	ctgagacaca	tgagactat	cttttgagaa	ctgggcaggt	tgtagatata	480
tcagacacca	tatacccgag	gaaccctgcc	atgtatagtg	aagaagctag	attaaagtcc	540
tttcagaact	ggccagacta	tgctcaccta	acccaagag	agttagcaag	tgctggactc	600
tactacacag	gtattgggtga	ccaagtgcag	tgcttttgtt	gtggtggaaa	actgaaaaat	660
tgggaacctt	gtgatcgtgc	ctggtcagaa	cacaggcgac	actttcctaa	tgcttctttt	720
gttttggggc	ggaatcttaa	tattcgaagt	gaatctgatg	ctgtgagttc	tgataggaat	780
ttcccaaatt	caacaaatct	tccaagaaat	ccatccatgg	cagattatga	agcacggatc	840
tttacttttg	ggacatggat	atactcagtt	aacaaggagc	agcttgcaag	agctggattt	900
tatgcttttag	gtgaagggtga	taaagtaaag	tgcttttact	gtggaggagg	gctaactgat	960
tggaagccca	gtgaagaccc	ttgggaacaa	catgctaaat	ggtatccagg	gtgcaaata	1020
ctgttagaac	agaagggaca	agaatatata	aacaatattc	atttaactca	ttcacttgag	1080
gagtgtctgg	taagaactac	tgagaaaaca	ccatcactaa	ctagaagaat	tgatgatacc	1140
atcttccaaa	atcctatgg	acaagaagct	atacgaatgg	ggttcagttt	caaggacatt	1200
aagaaaataa	tggaggaaaa	aattcagata	tctgggagca	actataaatc	acttgagggt	1260
ctggttgag	atctagtga	tgctcagaaa	gacagtatgc	aagatgagtc	aagtcagact	1320
tcattacaga	aagagattag	tactgaagag	cagctaaggc	gcctgcaaga	ggagaagctt	1380
tgcaaaatct	gtatggatag	aaatattgct	atcgtttttg	ttccttggtg	acatctagtc	1440
acttgtaaac	aatgtgctga	agcagttgac	aagtgtccca	tgtgctacac	agtcattact	1500
ttcaagcaaa	aaatttttat	gtcttaatct	aactctatag	taggcatggt	atgttggtct	1560
tattaccctg	attgaatgtg	tgatgtga				1588

<210> 4

<211> 236

<212> PRT

<213> Homo sapiens

<400> 4

Met	Ala	Asp	Tyr	Glu	Ala	Arg	Ile	Phe	Thr	Phe	Gly	Thr	Trp	Ile	Tyr
1				5				10						15	
Ser	Val	Asn	Lys	Glu	Gln	Leu	Ala	Arg	Ala	Gly	Phe	Tyr	Ala	Leu	Gly
		20						25					30		
Glu	Gly	Asp	Lys	Val	Lys	Cys	Phe	His	Cys	Gly	Gly	Gly	Leu	Thr	Asp
		35					40					45			
Trp	Lys	Pro	Ser	Glu	Asp	Pro	Trp	Glu	Gln	His	Ala	Lys	Trp	Tyr	Pro
	50					55					60				
Gly	Cys	Lys	Tyr	Leu	Leu	Glu	Gln	Lys	Gly	Gln	Glu	Tyr	Ile	Asn	Asn
65				70					75					80	
Ile	His	Leu	Thr	His	Ser	Leu	Glu	Glu	Cys	Leu	Val	Arg	Thr	Thr	Glu
			85					90					95		
Lys	Thr	Pro	Ser	Leu	Thr	Arg	Arg	Ile	Asp	Asp	Thr	Ile	Phe	Gln	Asn
		100						105					110		
Pro	Met	Val	Gln	Glu	Ala	Ile	Arg	Met	Gly	Phe	Ser	Phe	Lys	Asp	Ile
	115						120						125		
Lys	Lys	Ile	Met	Glu	Glu	Lys	Ile	Gln	Ile	Ser	Gly	Ser	Asn	Tyr	Lys
	130					135					140				

Ser Leu Glu Val Leu Val Ala Asp Leu Val Asn Ala Gln Lys Asp Ser
 145 150 155 160
 Met Gln Asp Glu Ser Ser Gln Thr Ser Leu Gln Lys Glu Ile Ser Thr
 165 170 175
 Glu Glu Gln Leu Arg Arg Leu Gln Glu Glu Lys Leu Cys Lys Ile Cys
 180 185 190
 Met Asp Arg Asn Ile Ala Ile Val Phe Val Pro Cys Gly His Leu Val
 195 200 205
 Thr Cys Lys Gln Cys Ala Glu Ala Val Asp Lys Cys Pro Met Cys Tyr
 210 215 220
 Thr Val Ile Thr Phe Lys Gln Lys Ile Phe Met Ser
 225 230 235

<210> 5
 <211> 500
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(500)
 <223> n = A,T,C or G

<400> 5
 caactacaca cgtgtgtgtg cgcgtgtgta taaaacacag tgcactaata ctcagccttt 60
 aaaaaaaatg ccacttgcaa caacgtagat ggagctggac gatatcatgc taaaattatg 120
 caaagtgaag caagcacaaa aaagaacgag acacgggagc ggggcacgag gtgctcactg 180
 ngcaagcgcc cactccaccg cgtgggttcc agctggagggc tgggagcggt ngtgggttcc 240
 tcttttcttg ctgacccttc ggagctctgg gaagtggctg caccttggcg gctccccaga 300
 gcgcgcgggt ctaatcgtag gtcgtcagcc tgggtggctg ggcccggtt agggcagggt 360
 ttggcatttc caatggtagg gggctcggac cgtccctccg cgggaccctc ccgttgggac 420
 aaggccgatc gcctgggagg ttggagccgc tctcctggcg cgagacgggt gacaagtcct 480
 atattcaaga gaagataact 500

<210> 6
 <211> 67
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Based on Cydia pomonella and Orgyia pseudotsugata

<221> VARIANT
 <222> (1)...(67)
 <223> Xaa = Any Amino Acid

<221> VARIANT
 <222> (1)...(67)
 <223> Xaa = Any Amino Acid

<400> 6

Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 35 40 45
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa Xaa Cys
 50 55 60
 Xaa Xaa Xaa
 65

<210> 7
 <211> 68
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Based on *Cydia pomonella* and *Orgyia pseudotsugata*

<221> VARIANT
 <222> (1)...(68)
 <223> Xaa = Any Amino Acid

<221> VARIANT
 <222> (1)...(68)
 <223> Xaa = Any Amino Acid

<400> 7
 Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 35 40 45
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa Xaa
 50 55 60
 Cys Xaa Xaa Xaa
 65

<210> 8
 <211> 69
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Based on *Cydia pomonella* and *Orgyia pseudotsugata*

<221> VARIANT
 <222> (1)...(69)
 <223> Xaa = Any Amino Acid

<221> VARIANT
 <222> (1)...(69)
 <223> Xaa = Any Amino Acid

<400> 8
 Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
 35 40 45
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa
 50 55 60
 Xaa Cys Xaa Xaa Xaa
 65

<210> 9
 <211> 70
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Based on Cydia pomonella and Orgyia pseudotsugata

<221> VARIANT
 <222> (1)...(70)
 <223> Xaa = Any Amino Acid

<221> VARIANT
 <222> (1)...(70)
 <223> Xaa = Any Amino Acid

<400> 9
 Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa
 35 40 45
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa
 50 55 60
 Xaa Xaa Cys Xaa Xaa Xaa
 65 70

<210> 10
 <211> 35
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Based on Cydia pomonella and Orgyia pseudotsugata

<221> VARIANT
<222> (1)...(35)
<223> Xaa = Any Amino Acid

<221> VARIANT
<222> (1)...(35)
<223> Xaa = Any Amino Acid

<400> 10
Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
1 5 10 15
Xaa His Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys
20 25 30
Xaa Xaa Cys
35